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Amendments to the Claims:

This listing of claims will replace all prior versions and listings of claims in the application:

Listing of Claims:

- 1 22. (Canceled).
- 23. (Currently amended) The polypeptide of claim 22-24 comprising a S. mutans competence signal peptide having SEQ ID NO: 4.
- 24. (Currently amended) A The polypeptide of claim 23 comprising all or part of an amino acid sequence in [SEQ ID NO:2] or [SEQ ID NO:4] and having S.mutans competence signal peptide activity.
- 25. (Currently amended) A polypeptide variant or a peptide mimetic of the polypeptide of claim 24SEQ ID NO: 2 or SEQ ID NO: 4 having at least 30% sequence identity to SEQ ID NO: 2 and having competence signal peptide activity or having activity for inhibiting the competence signal activity of the polypeptide of SEQ ID NO: 2 or 4.
- 26. (Currently amended) The polypeptide variant of claim 24 or 25 which is recombinantly produced.
- 27. (Currently amended) A polypeptide comprising a sequence The polypeptide variant of claim 25 having greater than 30%, 50% or 60% sequence identity to SEO ID NO: 2 or SEO ID NO: 4 the polypeptide of claim 24 or 25.
- 28. (Currently amended) The polypeptide variant of claim 24 or 25, isolated from S.mutans.
- 29-37. (Canceled).
- 38. (Currently amended) A-The polypeptide of claim 24 having an amino acid sequence wherein 1 15 amino acids of the polypeptide of claim 24 have been removed from the N- and/or COOH terminal of SEQ ID NO: 2 or SEQ ID NO: 4.

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- 39. (Currently amended) A—The polypeptide of claim 38 wherein 2 5 amino acids of the polypeptide of claim 24 have been removed from the N- and/or COOH terminal of SEQ ID NO: 2 or SEQ ID NO: 4.
- 40. (Currently amended) A-The polypeptide of claim 38 wherein 6 10 amino acids of the polypeptide of claim 24 have been removed from the N- and/or COOH terminal of SEQ ID NO: 2 or SEQ ID NO: 4.
- 41. (Currently amended) A-The polypeptide of claim 38 wherein 10 15 amino acids of the polypeptide of claim 24 have been removed from the N- and/or COOH terminal of SEQ ID NO: 2 or SEQ ID NO: 4.
- 42. (Currently amended) A synthetic polypeptide having competence signal peptide activity and comprising the 21-amino acid sequence of SEQ ID NO:164.
- 43. (Currently amended) A polypeptide having an amino acid sequence wherein 1 15 amino acids of the polypeptide of claim 24 amino acid sequence of SEQ ID NO: 2 or SEQ ID NO: 4 have been modified to include up to 1 point mutation per each 10 amino acids of SEQ ID NO: 2 or SEQ ID NO: 4, or a portion thereof.
- 44. (Currently amended) The polypeptide of claim 43 wherein each said mutation comprises comprises substitution with another amino acid.
- 45. (New) The polypeptide variant of claim 25 having greater than 60% sequence identity to SEQ ID NO: 2.
- 46. (New) The polypeptide variant of claim 45 having greater than 90% sequence identity to SEQ ID NO: 2.
- 47. (New) The polypeptide of claim 24 or the polypeptide variant of claim 25 having competence signal peptide activity.

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- 48. (New) The polypeptide of claim 24 or the polypeptide variant of claim 25 having activity for inhibiting the competence signal activity of the polypeptide of SEQ ID NO: 2 or 4.
- 49. (New) The polypeptide of claim 47 capable of binding to a bacterial histidine kinase cell surface receptor and activating said kinase.
- 50. (New) The polypeptide of claim 47 capable of promoting biofilm formation of said bacteria.
- 51. (New) The polypeptide of claim 47 capable of promoting acid tolerance in said bacteria.
- 52. (New) The polypeptide of claim 47 wherein said bacteria comprises Streptococcus mutans.
- 53. (New) The polypeptide of claim 48 capable of competitively inhibiting the binding of a peptide having SEQ ID NO. 4 to a bacterial histidine kinase cell surface receptor.
- 54. (New) The polypeptide of claim 48 capable of inhibiting biofilm formation of said bacteria.
- 55. (New) The polypeptide of claim 48 capable of inhibiting acid tolerance in said bacteria.
- 56. (New) The polypeptide of claim 48 wherein said bacteria comprises Streptococcus mutans.
- 57. (New) The polypeptide variant of any one of claims 25, 27, 45 and 46 comprising at least one modification chosen from the group consisting of
 - a N-terminal modification,
 - a COOH-terminal modification,
 - at least one amino acid is a D-amino acid, and
 - at least one chemically modified amino acid.